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<b>(54) Title:</b> RNA PACKAGING SYSTEM  <b>(57) Abstract</b> <p>The present invention relates to an <i>in vivo</i> system for expression and packaging of recombinant RNA into pseudovirus particles. The invention is based on the discovery that plant viral coat proteins (CPs) may be efficiently expressed in <i>E. coli</i>, and that these recombinant coat proteins will function to assemble <i>in vivo</i> and package recombinant chimeric RNA, containing an operatively linked origin-of-assembly (OAS) sequence, to form mature viral particles containing a foreign RNA. The present invention thus provides for packaging of RNA into a ribonuclease-resistant form that is easily purified and stored, and which overcomes the prior art problems associated with degradation of RNA by ribonucleases. Significantly, the method of the invention is RNA sequence- and length-independent. The components of the invention include a source in the bacterial host of viral coat proteins, and a source in the bacterial host to direct the transcription of a DNA molecule comprising an OAS-encoding DNA and a foreign DNA, which DNA molecule can be transcribed in the host cell to produce an RNA molecule comprising an OAS operatively linked to an RNA of interest. The CPs and OAS are from a plant virus having a rod-shaped helical particle and a single-stranded RNA genome, most preferably tobacco mosaic virus.</p>		

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## RNA PACKAGING SYSTEM

### 1. INTRODUCTION

The present invention relates to an in vivo  
5 system for expression and packaging of recombinant  
single-stranded RNA of unlimited length or sequence  
into pseudovirus particles comprised of plant viral  
coat proteins. The invention provides a means for  
efficient production and long-term handling and  
10 storage of otherwise labile RNA in ribonuclease-  
resistant virus-like particles.

### 2. BACKGROUND OF THE INVENTION

A number of infectious plant viruses,  
15 including members of the Tobamo-, Potex-, Poty- and  
Tobra-groups of viruses, share properties in common  
with one another. These properties include a single-  
stranded RNA genome encapsidated by viral coat protein  
oligomers that assemble to form either elongated rigid  
20 rods or flexuous threads.

Perhaps the best studied of the plant  
viruses is the tobamovirus Tobacco Mosaic Virus (TMV)  
which has a genome size of 6.4 Kb. The positive  
stranded genomic RNA codes for a number of viral  
25 proteins including those required for replication of  
the viral genome and those coding for structural  
proteins such as the coat protein which assembles into  
20S protohelical or disk-like structures that become  
arranged into elongated helical structures with the  
30 viral genomic RNA molecule (Goelet et al., 1982, Proc.  
Natl. Acad. Sci. 79:5818-22).

Contiguous with the TMV genomic RNA is a  
sequence element referred to as the origin-of-assembly  
sequence (OAS) that is necessary and sufficient to  
35 direct efficient encapsidation of contiguous viral RNA

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sequences into virus particles. The TMV OAS is located approximately 1 Kb from the 3' end of the viral genome in the common strain (and in the coat protein gene itself in the cowpea strain (Cc; Sunn-hemp mosaic virus)) and consists of a 440 nucleotide sequence that is predicted to form three hairpin stem-loop structures (Turner and Butler, 1986, Nucl. Acids Res. 14:9229-42). The viral coat protein disks initially bind to loop 1 (the 3' most) during viral assembly and *in vitro* packaging assays using mutant assembly origin transcripts have defined the 75 nucleotides comprising loop 1 as necessary and sufficient for encapsidation of foreign or viral RNA sequences (Turner et al., 1988, J. Mol. Biol. 203:531-47).

*In vitro* reconstitution studies have provided details on the assembly process for TMV. Preparations of purified coat protein, derived from virions from infected plant cells, are able to assemble into helical structures and virus-like rods, even in the absence of RNA at pH 5, suggesting that the coat protein contains the essential information required for self-assembly. Incubation of purified TMV coat protein preparations with TMV RNA at pH 7, *in vitro*, results in assembly of TMV-like viral particles containing encapsidated RNA (Fraenkel-Conrat and Williams, 1955, Proc. Natl. Acad. Sci. 41:690-98). Furthermore, it has been shown that foreign chimeric RNA molecules containing OAS sequences, transcribed *in vitro* using SP6 or T7 (Jupin, I. et al., 1989, Nucl. Acids Res. 17:815) transcription plasmids can also be assembled *in vitro* into pseudovirus particles (Sleat et al., 1986, Virology 155:299-308).

Until recently, sources of viral coat proteins for *in vitro* reconstitution studies have

relied on virus preparations made from infected plant tissue. However, such sources are disadvantageous since laborious procedures comprising infection with virus, purification of the virus from plant tissue, and then purification of coat protein from the virus must be used. The cloning and sequencing of a number of plant viral genomes has led to the identification of viral coat protein encoding sequences. Insertion of these genes into bacterial expression vectors has allowed the expression of, for instance, TMV coat protein in *E. coli* (Shire et al., 1990, *Biochemistry* 29:5119-26). However, it was reported (*id.*) that recombinant TMV coat protein produced in *E. coli* reconstitutes *in vitro* with TMV RNA at a greatly reduced rate relative to the reconstitution with native coat protein; the authors suggested that this inefficiency in reconstitution arises from the lack of an acetyl group on the amino terminus of the recombinant protein, which is present on the native coat protein. Zucchini yellow mosaic virus and Johnsongrass mosaic virus (both potyviruses, which are not members of the tobamovirus, tobnavirus, or potexvirus groups) coat proteins have also been produced in *E. coli* (Gal-On et al., 1990, *Gene* 87:273-277; Jagadish et al. 1991, *J. Gen. Virology* 72:1543-1550).

Present work in molecular biology and recombinant nucleic acid technology is encumbered by problems associated with degradation of RNA by ribonucleases. Researchers in the past have relied on inhibitors such as human placental RNase inhibitor (RNasin), or the use of alkylating reagents such as diethylpyrocarbonate (DEPC) which is a suspected carcinogen to inhibit the activity of ribonucleases;

such inhibitors may produce (DEPC) undesirable modified components of RNA.

### 3. SUMMARY OF THE INVENTION

5           The present invention relates to an *in vivo* system for expression and packaging of recombinant single-stranded RNA into pseudovirus particles. The invention is based on the discovery that viral coat proteins (CPs) may be efficiently expressed in *E.*  
10 *coli*, and that these recombinant coat proteins will function to assemble *in vivo* and package recombinant chimeric RNA, containing a contiguous origin-of-assembly sequence (OAS), to form mature virus-like particles containing a foreign RNA. The present  
15 invention thus provides for packaging of RNA into a ribonuclease-resistant form that is easily purified and stored, and which overcomes the prior art problems associated with degradation of RNA by ribonucleases. The invention provides for convenient, efficient  
20 production and long-term storage of any RNA of interest, without substantial degradation. Significantly, the method of the invention is RNA sequence- and length-independent. The *in vivo* system provided herein for production and packaging of  
25 recombinant RNA will have broad applications that will include any of the techniques currently used by molecular biologists involving manipulation of RNA. These include for example *in vitro* translation reactions to produce proteins of interest, synthesis  
30 of sense and anti-sense RNA molecules for introduction into cells to study protein function, and synthesis of radiolabelled RNA probes for use in Northern or Southern Blot analyses."

          The components of the invention include a  
35 source in the bacterial host of viral coat proteins,

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and a source in the bacterial host to direct the transcription of a DNA molecule comprising an OAS encoding DNA linked to a foreign DNA (hereinafter "\*DNA"), which DNA molecule can be transcribed in the host cell to produce an RNA molecule comprising an OAS linked to an RNA of interest (hereinafter "OAS-linked \*RNA"); both sources being compatible and capable of protein (CP) expression and transcription (of OAS-linked \*DNA), respectively, together in the same bacterial host. The CPs and OAS provided by the invention can be that of any plant virus having a rod-shaped helical particle and a single-stranded RNA genome, including but not limited to a tobamovirus, potexvirus, tobnavirus, hordeivirus, potyvirus, and furovirus. In a preferred aspect, the sources of the CP and OAS-linked \*DNA are plasmid vectors. In a specific embodiment, a single plasmid vector is the source of both CP and OAS-linked \*RNA. In another embodiment, the different sources are different plasmid vectors. Vectors, recombinant cells, and kits for carrying out the method of the invention are also provided.

#### 4. DESCRIPTION OF THE FIGURES

Figure 1. Coomassie stained (15%) polyacrylamide gel of TMV coat proteins expressed in *E. coli*. Lanes: 1, pETPro301 (+) IPTG; 2, pET302 (+) IPTG; 3, pETAla301 (+) IPTG; 4, pET301 (+) IPTG; 5, pET3a (+) IPTG; 6, pET3a (-) IPTG; 7, protein size markers.

Figure 2. Western blot analysis of TMV coat proteins expressed in *E. coli*. Lanes: 1, pETAla301 (+) IPTG; 2, pET301 (+) IPTG; 3, pET301 (-) IPTG; 4, pET302 (+) IPTG.

Figure 3. Electron micrographs of *E. coli*-produced TMV coat protein and TMV-like particles.



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Immunosorbent electron microscopy of complexes containing TMV coat protein from IPTG-induced cultures of *E. coli* transformed with pETAla301 and pLysE (negative control with CP alone; panel A), or

5 pETAla301 and pLys102 (panels B, C, D; = CAT-OAS mRNA) was performed. Negative staining was with 1% uranyl acetate (panels A, B, C) or with 1% ammonium molybdate (panel D). Panel E shows a histogram of TMV-like particles.

10 Figure 4. Northern dot blots of RNA extracted from virus-like particles. Lanes: 1, H<sub>2</sub>O control; 2, pETAla301 and pLys102 (CAT-OAS mRNA); 3, pETAla301 and pLys1032 (GUS-OAS mRNA); 4, pLysE and pETAla301.

Figure 5. PCR amplification of RNA samples  
15 isolated from pseudovirus particles by use of 5' and 3' primers derived from the TMV OAS sequence. Lanes: 1, TMV RNA (positive control); 2, molecular weight "markers"; 3, pLysE CAT; 4, pLys102 CAT-OAS; 5, pLys1032 GUS-OAS.

20 Figure 6. Schematic diagram of a portion of the pET3a vector derivatives encoding TMV CP. The thin line represents the pET3a vector sequence. The site directed mutagenesis of the 5' end of the gene encoding TMV CP was done by PCR reactions. The  
25 mutated gene encoding TMV CP was cloned into the final expression vector, pET3a, using the NdeI and BamHI sites.

Figure 7. Vector providing OAS-linked \*DNA. A schematic diagram of a portion of the pLys102 vector  
30 providing the transcript to be packaged in *E. coli*, comprising the OAS-linked CAT DNA, is shown. OAS is the origin of assembly sequence of TMV. The hatched area shows a repeated sequence of the CAT gene generated by cloning-in the OAS cassette.

35

Figure 8: Vector providing OAS-linked \*DNA. A schematic diagram of a portion of the pLys1032 vector providing the transcript to be packaged in *E. coli*, comprising the OAS-linked GUS DNA, is shown. OAS shows the origin of assembly sequence of TMV. The hatched area shows the synthetic poly (CAA) leader.

Figure 9. Schematic diagram of OAS probe. A schematic diagram is shown of a portion of pJII102 (Gallie et al., 1987, Science 236:1122-24) comprising the sequence from which the OAS hybridization probe was derived.  $\Omega'$  represents TMV leader derivative (Gallie et al., 1987, Nucl. Acids Res. 15:3127).

Figure 10. TMV origin of assembly. The nucleotide sequence from nucleotides 5118-5550 (SEQ ID NO:5) of the TMV genome (from Goelet et al., 1982, Proc. Natl. Acad. Sci. USA 79:5818-22), containing the TMV OAS extending from bases 5290 to 5527 of the viral sequence (nucleotides 173-410 in SEQ ID NO:5) (Zimmern, D., 1983, EMBO J. 2:1901-07), is shown.

Figure 11. Loop 1 of TMV OAS. The nucleotide sequence (SEQ ID NO:6) is shown of loop 1 of the TMV OAS, which is known to be sufficient to direct the encapsidation of a contiguous RNA fragment by TMV coat protein disks (from Turner, D.R., et al., 1988, J. Mol. Biol. 203:531-47; Fig. 1).

## 5. DETAILED DESCRIPTION OF THE INVENTION

The present invention relates to an *in vivo* system for production and packaging of recombinant single-stranded RNA into pseudovirus (virus-like) particles. The *in vivo* packaging system is useful as a means of producing significant amounts of RNA of interest in a form that is both easily purified and protected from the degradative effects of ribonucleases.

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The invention is based on the ability to express, *in vivo*, viral coat proteins (CPs) and package chimeric recombinant RNA containing OAS-linked sequences (hereinafter "OAS-linked \*RNA") such that

5 the CPs function to assemble and package the OAS-linked \*RNA into pseudovirus particles. As will be apparent, "OAS-linked \*DNA" refers herein to a DNA molecule comprising a sequence which can be transcribed to an OAS RNA sequence linked to an RNA of

10 interest, while "OAS-linked \*RNA" refers to the RNA sequence thus produced by such transcription. The components of the invention include sources for production within a bacterial host of both the viral CPs and the OAS-linked \*RNA. In one aspect, the

15 recombinant CP gene may be integrated into a bacterial host chromosome (*e.g.*, by homologous recombination; see, *e.g.*, Koller and Smithies, 1989, *Proc. Natl. Acad. Sci.* 86:8932-35; Zijlstra et al., 1989, *Nature* 342:435-38) for expression therefrom, and the

20 OAS-linked \*DNA construct may be integrated into a bacterial host chromosome (*e.g.*, by homologous recombination) for transcription therefrom. In an alternative, preferred embodiment, the CP is expressed from a vector, preferably a plasmid expression vector,

25 and the OAS-linked \*DNA is transcribed from a vector, preferably a plasmid expression vector, which can be the same or a different vector from the expression vector encoding the CP. Sources of the CP and OAS-linked \*RNA according to the invention are detailed

30 below.

### 5.1. PRODUCTION OF VIRAL COAT PROTEIN IN A BACTERIAL HOST

A first component of the invention is a source providing for recombinant production in a bacterial host of a plant viral CP. In a preferred aspect, the CP nucleic acid encodes a CP of a virus that is the same virus strain from which the OAS is derived.

The plant viral CP is the CP of a plant virus having a rod-shaped helical particle and a single-stranded RNA genome, including but not limited to a tobamovirus, potexvirus, tobnavirus, hordeivirus, potyvirus, and furovirus, with the tobamovirus TMV most preferred. These virus groups have rod-shaped helical particles of lengths determined only by the size of the RNA therein; i.e., there are no packaging size constraints. Both natural viral strains and nitrous acid- and other induced mutant strains are contemplated as included in the foregoing virus groups. Members of the tobamovirus group include but are not limited to TMV, cucumber green mottle mosaic virus, tomato mosaic virus, Frangipani mosaic virus, Odontoglossum ringspot virus, Holmes' Ribgrass mosaic virus, sammons' Opuntia virus, Sunn-hemp mosaic virus (cowpea Cc), U2-tobacco mosaic virus, etc. (see, e.g., CMI/AAB Descriptions of Plant Viruses, September 1977, No. 184). Tobnaviruses include but are not limited to pea early-browning virus, tobacco rattle virus, and pepper ringspot virus. Potexviruses include but are not limited to potato virus X and papaya mosaic virus. Hordeiviruses include but are not limited to barley stripe mosaic virus. Potyviruses include but are not limited to potato virus-Y. Furoviruses include but are not limited to soil-borne wheat mosaic virus, beet necrotic yellow vein virus, and potato mop-top virus.

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The nucleic acid encoding the CP is obtained by any means available in the art. Various CP nucleic acid sequences have been disclosed and can be used. For example, the TMV CP has been cloned and sequenced (see Goelet et al., 1982, Proc. Natl. Acad. Sci. 79:5818-22). Additionally, the sequence of a TMV CP gene with codons optimized for translation in a prokaryotic system has been disclosed (Haynes et al., 1986, Biotechnology 4:637-41). If a nucleic acid clone of the desired CP is not already available, the clone can be obtained by use of standard recombinant DNA methodology. For example, the DNA may be obtained by standard procedures known in the art from cloned DNA (e.g., a DNA "library"), by chemical synthesis, or by isolation of the viral RNA or fragments thereof, purified, preferably, from viral particles (see, for example Sambrook et al., 1989, Molecular Cloning, A Laboratory Manual, 2nd Ed., Cold Spring Harbor Laboratory, Cold Spring Harbor, New York; Glover, D.M. (ed.), 1985, DNA Cloning: A Practical Approach, MRL Press, Ltd., Oxford, U.K. Vol. I, II.)

Identification of the specific DNA fragment containing the desired gene may be accomplished in a number of ways. For example, if an amount of a portion of a CP DNA or its specific RNA, or a fragment thereof, is available and can be purified and labeled, the generated DNA or RNA may be screened by nucleic acid hybridization to the labeled probe (Benton, W. and Davis, R., 1977, Science 196:180; Grunstein, M. and Hogness, D., 1975, Proc. Natl. Acad. Sci. U.S.A. 72:3961). Those DNA fragments with substantial homology to the probe will hybridize. It is also possible to identify the appropriate DNA by restriction enzyme digestion(s) and comparison of fragment sizes with those expected according to a

known restriction map if such is available. Further selection can be carried out on the basis of the properties of the gene. Alternatively, the presence of the gene may be detected by assays based on the physical, chemical, or immunological properties of its expressed product. For example, a cDNA clone can be selected which produces a CP that, e.g., has similar or identical electrophoretic migration, isoelectric focusing behavior, proteolytic digestion maps, self-assembly activity or antigenic properties as known for the purified CP.

An appropriate CP RNA can also be identified by *in vitro* translation. Immunoprecipitation analysis or functional assays (e.g., self-assembly ability in *vitro*) of the *in vitro* translation products of the isolated RNA identifies the RNA as one containing the desired sequences.

Alternatives to isolating a CP gene include, but are not limited to, chemically synthesizing the gene sequence itself from a known sequence or making cDNA to the RNA which encodes the CP. Other methods are possible and within the scope of the invention.

If desired, the identified and isolated gene can then optimally be inserted into an appropriate cloning vector prior to transfer to a bacterial expression vector of the invention.

Nucleic acids which encode derivatives (including fragments) and analogs of native CPs can also be used in the present invention, as long as such derivatives and analogs retain the ability to assemble into a viral particle and package therein an OAS-containing RNA. In particular, CP derivatives can be made by altering CP sequences by substitutions, additions, or deletions that provide for functionally active molecules. Furthermore, due to the inherent

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degeneracy of nucleotide coding sequences, other DNA sequences which encode substantially the same or a functionally equivalent viral CP amino acid sequence may be used in the practice of the methods of the invention. For example, it may be useful to change DNA sequences to optimize for *E. coli* or other bacteria codon usage, or changes may be made in amino acid sequences to enhance viral particle assembly in the bacterial host system. Such alterations of the coat nucleotide sequence include deletions, additions or substitutions of different nucleotides resulting in a sequence that encodes the same or a functionally equivalent gene product. The gene product may contain deletions, additions or substitutions of amino acid residues within the sequence which result in silent changes thus producing a bioactive product. Such amino acid substitutions may be made on the basis of similarity in polarity, charge, solubility, hydrophobicity, hydrophilicity and/or the amphipathic nature of the residues involved. For example, negatively charged amino acids include aspartic acid and glutamic acid; positively charged amino acids include lysine and arginine; amino acids with uncharged polar head groups or nonpolar head groups having similar hydrophilicity values include the following: leucine, isoleucine, valine; glycine, alanine; asparagine, glutamine; serine, threonine; phenylalanine, tyrosine.

In a specific embodiment, nucleic acids can be made which encode a CP fusion protein derivative which comprises a surface ligand fused via a peptide bond to the CP, such that the presence of the surface ligand does not interfere with OAS-dependent RNA assembly. "Surface ligand" as used herein refers to a peptide or protein which binds to a receptor on the

surface of a cell. Thus, such surface ligand-CP fusion proteins can target the pseudovirus particle *in vivo* to a particular cell type expressing the receptor for the surface ligand; subsequent receptor-mediated  
5 endocytosis can effect intracellular delivery of the particle and its contained RNA into the cell. Surface ligands which can be used include but are not limited to peptides containing the Arg-Gly-Asp amino acid motif (the ligand for the polio receptor),  
10 asialoglycoprotein, etc.

Derivatives or analogs of CPs also include but are not limited to those peptides which are substantially homologous to CP or fragments thereof, or whose encoding nucleic acid is capable of  
15 hybridizing to a CP nucleic acid sequence, which are functionally active in assembly and packaging.

The nucleic acids encoding derivatives and analogs of the CP can be produced by various methods known in the art. For example, the cloned CP gene  
20 sequence can be modified by any of numerous strategies known in the art (Sambrook et al., 1989, Molecular Cloning, A Laboratory Manual, 2nd ed., Cold Spring Harbor Laboratory, Cold Spring Harbor, New York). The sequence can be cleaved at appropriate sites with  
25 restriction endonuclease(s), followed by further enzymatic modification if desired, isolated, and ligated *in vitro*. In the production of the gene encoding a derivative or analog of a CP, care should be taken to ensure that the modified gene remains  
30 within the same translational reading frame as the CP, uninterrupted by translational stop signals, in the gene region where the CP functional activity is encoded.

Additionally, the CP-encoding nucleic acid  
35 sequence can be mutated *in vitro* or *in vivo*, to create



and/or destroy translation, initiation, and/or termination sequences, or to create variations in coding regions and/or form new restriction endonuclease sites or destroy pre-existing ones, to  
5 facilitate further *in vitro* modification. Any technique for mutagenesis known in the art can be used, including but not limited to, *in vitro* site-directed mutagenesis (Hutchinson, C., et al., 1978, J. Biol. Chem 253:6551).

10 The desired CP-encoding nucleic acid is then preferably inserted into an appropriate bacterial expression vector, i.e., a vector which contains the necessary elements for the transcription and  
15 translation of the inserted protein-coding sequence in a bacterial host, to create a vector that functions to direct the synthesis of the CP that will subsequently assemble and encapsidate the OAS-linked \*RNA within the bacterium. A variety of vector systems may be utilized to express the CP-coding sequence, but the  
20 vector must be functional in the bacterial host and compatible with any other vector present (e.g., a vector containing the OAS-linked \*DNA). Such vectors include but are not limited to bacteriophages, plasmids, or cosmids. In a preferred aspect, a  
25 plasmid expression vector is used. The expression elements of vectors vary in their strengths and specificities. Any one of a number of suitable transcription and translation elements may be used, as long as they are functional in the bacterial host.

30 Standard recombinant DNA methods may be used to construct expression vectors containing a chimeric gene consisting of appropriate transcriptional/translational control signals and the CP coding sequences (see, e.g., Sambrook et al., 1989,  
35 *supra*). These methods may include *in vitro*

recombinant DNA and synthetic techniques and *in vivo* recombinants (genetic recombination). The insertion into a cloning vector can, for example, be accomplished by ligating the DNA fragment into a cloning vector which has complementary cohesive termini. However, if the complementary restriction sites used to fragment the DNA are not present in the cloning vector, the ends of the DNA molecules may be enzymatically modified. Alternatively, any site desired may be produced by ligating nucleotide sequences (linkers) onto the DNA termini; these ligated linkers may comprise specific chemically synthesized oligonucleotides encoding restriction endonuclease recognition sequences. In an alternative method, the cleaved vector and CP gene may be modified by homopolymeric tailing. Expression of the nucleic acid sequence encoding the CP may be regulated by a second nucleic acid sequence. For example, expression of a CP may be controlled by any bacterial promoter/enhancer element known in the art. Promoters which may be used to control CP gene expression can be constitutive or inducible, and include, but are not limited to, the  $\beta$ -lactamase promoter (Villa-Komaroff, et al., 1978, Proc. Natl. Acad. Sci. 75:3727-31), the tac promoter (DeBoer, et al., 1983, Proc. Natl. Acad. Sci. 80:21-25), lac UV5,  $\lambda P_1$ , T7, and trp promoters.

Alternatives to isolating a CP gene include, but are not limited to, chemically synthesizing the gene sequence itself from a known sequence or making cDNA to the RNA which encodes the CP gene. Other methods are possible and within the scope of the invention.

In addition to promoter sequences, the CP expression vectors preferably contain specific initiation signals for efficient translation of

inserted coat protein sequences. One such signal, referred to as the Shine-Dalgarno sequence, acts as a ribosome binding site that is necessary for efficient translation of RNA. In *E. coli*, the ribosome binding site includes an initiation codon (ATG) and a sequence of 3-9 nucleotides in length located 3-11 nucleotides upstream from the initiation codon (e.g., of the CP gene); on average, six out of eight nucleotides in the mRNA match this sequence: 5'-UAAGGAGG-3'.

10 Transcription termination signals (downstream of the CP gene), selectable markers such as genes encoding antibiotic resistance, and a gene providing for expression of lysozyme (which, *inter alia*, aids in later purification of the pseudovirus particle by

15 making the bacterial cell walls less resistant to rupture) are preferably also included in the plasmid expression vectors. In the BH21 (DE3) system T7 lysozyme is apparently also a stringent control for

20 gene for the T7 RNA polymerase, since it functions to bind T7 polymerase.

In a specific embodiment, a recombinant expression vector provided by the invention, encoding a plant virus CP or functional derivative thereof,

25 comprises the following, operatively linked components: a promoter which controls the expression of the CP or functional derivative thereof, a translation initiation signal, a DNA sequence encoding the CP or functional derivative thereof, and a

30 transcription termination signal. In a preferred aspect, the above components are present in 5' to 3' order as listed above.

In another specific embodiment, described in the examples sections herein, the gene encoding the

35 TMV CP is inserted downstream of both the T7 promoter

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from phage T7 gene 10 and the translational initiation signal (Shine-Dalgarno site) from gene 10 of this phage. As described *infra*, both wild type CP gene sequences (plasmid pET301) and CP sequences optimizing  
5 for *E. coli* codon usage (plasmid pET302) were used, as were CPs with changes in the penultimate amino-terminal Ser (the N-formyl methionine is removed by post-translational processing) to either Ala, Pro, or Asp. A T7 RNA polymerase terminator and a *colE1*  
10 replicon were also included in the expression vector. In this embodiment, T7 RNA polymerase transcribes the CP sequence. The T7 RNA polymerase is produced from a chromosomally integrated sequence (a  $\lambda$  derivative lysogen) under the control of an inducible promoter,  
15 lac UV5. Thus, when lac UV5 transcription is induced by IPTG, T7 RNA polymerase is produced, which in turn transcribes the CP gene to produce CP. Use of an expression vector containing the T7 promoter is preferred since it has a number of advantages  
20 contributing to efficient transcription of genes cloned adjacent to the T7 promoter. These include the strict specificity of T7 polymerase for its cognate promoter sequences, the rapid rate at which elongation proceeds and the infrequency at which premature  
25 termination of transcription occurs.

#### 5.2. PRODUCTION OF OAS-LINKED \*RNA IN THE BACTERIAL HOST

A second component of the invention is a  
30 source providing for transcription of the OAS-linked \*DNA, in the same bacterial host in which the CP is expressed, whereby the expressed CP assembles into a viral particle and packages therein the OAS-linked \*RNA resulting from transcription.

35 The viral origin-of-assembly sequences (OAS) are an essential feature of the OAS-linked \*DNA

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vectors as they are responsible for directing the encapsidation of the heterologous RNA molecules (the OAS-linked \*RNA) into pseudovirus particles. OAS, as used herein, refers to an RNA sequence (or DNA

5 encoding such an RNA sequence) that can be specifically recognized by plant viral CPs, and which is sufficient to direct the encapsidation by such CPs of an RNA containing the OAS. As will be apparent, the OAS need not be directly linked at the 5' or 3'

10 end of the \*DNA; for example, there may be some intervening sequence; what is required is that the OAS be positioned with regard to the \*DNA so that, upon transcription, a ribonucleic acid is produced (OAS-linked \*RNA) wherein the OAS is operatively

15 linked to the RNA sequences of interest so as to enable the encapsidation of the RNA molecule into virus-like particles by the CP. For example, transcription termination signals should only occur downstream of the OAS-linked \*DNA (transcription

20 proceeding in an upstream to downstream direction).

The OAS-linked \*DNA is preferably inserted in an appropriate vector, preferably a plasmid vector, which can direct the transcription within the bacterium of OAS-linked \*DNA to form the OAS-linked

25 \*RNA. The choice of vector, and its other components (e.g., promoter, selectable marker, etc.) can be as described *supra* in Section 5.1, but the vector must have regulatory elements functional in and compatible with the bacterial host, and compatible with the CP

30 expression vector also present in the host. Such an OAS-linked \*DNA vector preferably comprises the following operatively linked components: promoter, OAS, \*DNA, and transcriptional termination signal. In one embodiment, the foregoing components are present

35 in 5' to 3' order in which they are listed. In an

alternative embodiment, the OAS-linked \*DNA vector comprises, in 5' to 3' order: promoter, \*DNA, OAS, and transcriptional termination signal. This latter embodiment takes advantage of the fact that an OAS (e.g., the TMV OAS) can function in a bidirectional manner and can direct encapsidation when located at either the 5' or 3' end of the RNA; however, positioning the OAS at the 5' end of the \*DNA is preferred in order to inhibit degradation of its transcribed RNA within the cell. Thus, in designing the OAS-linked \*DNA expression vectors it may be most effective to place the OAS sequences at the 5' end of the \*DNA so that newly transcribed RNA can be rapidly and efficiently assembled into viral particles. In this way, co-transcriptional encapsidation should protect the RNA with coat protein as the RNA is being synthesized, thereby circumventing problems associated with rapid RNA degradation *in vivo*. However, in an embodiment where the \*RNA is an mRNA to be translated into protein, a 5'-OAS sequence may interfere with translation of the mRNA, unless an IRES or Shine-Dalgarno 40S/30S ribosome binding site is engineered between the OAS and \*DNA. Alternatively, a self-cleavage ribozyme site may be inserted between the OAS and DNA.

In a specific embodiment, the OAS-linked \*DNA vector further comprises a polylinker region, comprising a variety of restriction endonuclease cleavage sites, situated just 5' or 3' to the \*DNA.

In a preferred aspect, the OAS-linked \*DNA expression vector also contains a selectable marker gene, such as one encoding antibiotic (tetracycline, chloramphenicol, neomycin, etc) resistance.

In a preferred aspect, the RNA component of the OAS-linked \*RNA is a mRNA encoding a protein of interest, and thus, preferably, to facilitate eventual

translation thereof (see Section 5.4) a Shine-Dalgarno sequence is incorporated before the coding sequence in the \*DNA.

The vectors may optionally and preferably also provide for expression of lysozyme. In a specific aspect, when using the T7 promoter system it may also be useful to express the T7 lysozyme protein, since it functions to bind T7 polymerase and inhibit low level constitutive expression by T7 polymerase. It is also preferred to express T7 lysozyme since this results in bacterial cells that are more fragile due to degradation of the *E. coli* peptidoglycan wall by lysozyme. This feature becomes useful at later stages when the bacterial cells are lysed as a step in the purification of viral particles. The T7 lysozyme gene may be expressed in the bacterial host cell either chromosomally or episomally.

As discussed supra for the CP expression vectors, methods which are well known to those skilled in the art can be used to construct bacterial vectors containing OAS-linked \*DNA and appropriate transcriptional control sequences. These methods include *in vitro* recombinant DNA techniques (see, for example, Sambrook et al., 1989, Molecular Cloning, A Laboratory Manual, 2nd. Ed., Cold Spring Harbor Laboratory, N.Y).

In a specific embodiment, the promoter directing the transcription of the OAS-linked \*DNA is inducible and is the same promoter as that which, present on a different plasmid, directs CP gene expression. Alternatively, the CP gene promoter may be a different inducible promoter, or may be a constitutive promoter, or the OAS-linked \*DNA promoter may be constitutive. As will be apparent, any combination of promoter types may be used.

One of the advantages of the production and packaging system provided by the present invention is that expression and packaging appear to be both sequence- and length-independent with respect to the

5 \*RNA component of the OAS-linked \*RNA. As a result, any DNA sequence of interest may be linked to an OAS and then transcribed and its RNA assembled into virus-like particles according to the invention. Thus, in a specific embodiment of the invention, a plasmid vector

10 is provided comprising the following operatively linked elements: bacterial promoter, polylinker, OAS, transcriptional termination signal. In one embodiment, the foregoing elements appear in 5' to 3' order in which they are listed above. In another

15 embodiment, the vector comprises the following, listed in 5' to 3' order: bacterial promoter, OAS, polylinker, transcriptional termination signal. Such vectors facilitate the insertion of any DNA sequences of interest, by virtue of the polylinker region

20 containing a variety of restriction endonuclease cleavage sites into which the foreign \*DNA may be inserted.

The selection of specific plant viral OAS sequences for use will depend on what plant viral CPs

25 are being expressed in the bacterial host cell. It is important to keep in mind that the viral OAS sequences and viral CPs must be functionally compatible with one another for efficient assembly of OAS-linked \*RNA into virus particles, i.e., the CP expressed must be able

30 to undergo particle assembly and to recognize the OAS for encapsidation therein. OAS sequences which can be used thus include but are not limited to those of rod-shaped plant viruses including the tobnaviruses, tobamoviruses, and potexviruses, with a TMV OAS most

35



preferred. In a preferred aspect, the OAS and CP sequences are derived from the same strain of virus.

Perhaps the most well-defined of the origin-of-assembly sequences is that comprising the TMV OAS  
5 (see, e.g., Turner et al., 1988, J. Mol. Biol. 203(3):531-47; Turner and Butler, 1986, Nucl. Acids Res. 14(23):9229-42). Experiments with TMV have defined an approximately 440 nucleotide sequence, located 1 Kb from the 3' end of viral genomic RNA,  
10 that comprises the OAS sequence (Figure 10; SEQ ID NO:5). This region is found to contain three regions predicted to form three stable hairpin stem-loops (Zimmern, D., 1983, EMBO J. 2:1901-07), and it has been shown that coat protein 20S "disks" initially  
15 bind to loop 1 (the 3' most) during viral assembly. It has also been demonstrated using *in vitro* packaging assays and mutant assembly origin transcripts (Turner, D.R., et al., 1988, J. Mol. Biol. 203:531-47), that rapid and specific assembly initiation may occur in  
20 the absence of loop 2 and 3, but that loop 1 sequences, which consist of about 75 nucleotides (Fig. 11; SEQ ID NO:6) are necessary and sufficient for encapsidation of heterologous RNA sequence. It is these 75 nucleotides that are the preferred OAS  
25 sequences for use in construction of OAS-linked \*DNA expression vectors.

In other embodiments of the invention, an OAS other than the TMV OAS discussed above can be used. For example, cowpea strain TMV has an OAS in  
30 the middle of its CP gene (Meshi et al., 1981, Mol. Gen. Genet. 184:20-25) which can be used; the CP gene (containing the OAS) can thus be ligated to a \*DNA to form an OAS-linked \*DNA. In this embodiment, the OAS-linked \*DNA would also function as the CP source,  
35 obviating the need for an additional source of CP. In

another specific embodiment, the OAS at the 5' end of the potexvirus papaya mosaic virus RNA (Lok and Abou-Haidar, 1986, Virology 153:289-96) can be used.

In a preferred aspect of the invention, the  
5 OAS-linked \*RNA and CP are produced from different expression vectors within the bacterial host. However, in an alternative embodiment, the CP gene can be incorporated in and expressed from the vector also producing the OAS-linked \*RNA.

10 In specific embodiments of the invention detailed in the examples sections, a two-part approach is used to generate OAS-linked \*RNA: Firstly, T7 RNA polymerase is expressed under the control of an inducible promoter such as lac UV5; secondly, a  
15 plasmid vector expresses the OAS-linked \*DNA under the control of a T7 RNA promoter. Thus, when T7 RNA polymerase is expressed upon induction by, e.g., IPTG, the expressed T7 RNA polymerase will recognize the T7 promoter and transcribe the OAS-linked \*DNA. In the  
20 specific embodiment detailed in the examples section, the OAS-linked \*DNA vector contained the T7 promoter followed by the  $\beta$ -glucuronidase (GUS) gene cassette. The TMV OAS sequence and a T7 terminator were located at the 3' side of the GUS coding sequence. As  
25 detailed *infra*, a third expression vector supplied the CP for encapsidation, also expressed under control of the T7 promoter.

In another specific embodiment of the invention detailed in the examples *infra*, OAS-linked  
30 \*RNA was produced from a plasmid containing the chloramphenicol acetyl transferase (CAT or Cm') gene followed by the TMV OAS. Low level constitutive transcription was from the *E. coli* promoter of the Cm' gene.

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5.2.1. BACTERIAL HOSTS

Although *E. coli* is the preferred host organism, other strains of bacteria including but not limited to *Salmonella* spp., *B. subtilis*, and transformable *Cyanobacteria* (*Synechococcus*, *Anabena*, *Anacystis nidulans*) may be used for expression of OAS-linked \*RNA and CP. The choice of bacterial strains will depend on which expression vectors (whether episomal or chromosomally integrated) are used for expression of viral CP and to direct the transcription of OAS-linked \*DNA, since the regulatory elements of the vectors must be functional in and compatible with the bacterial host and any other vectors present. For instance, promoters, transcriptional termination signals, and origins of replication (for plasmids) should be chosen that are functional in the bacterial strain chosen, based on knowledge common in the art or readily available to the skilled artisan. As another example, if the expression of OAS-linked \*DNA is under the control of the  $\lambda P_L$  promoter, the cIts857 repressor protein should be supplied by the host organism (or by expression of a recombinant plasmid therein). Similarly, if expression is controlled by a T7 promoter, a source of T7 RNA polymerase should be available in the host cell. As stated *supra*, the host bacterium may contain the CP DNA sequence and OAS-linked \*DNA constructs both episomally or both chromosomally integrated or one episomally and one chromosomally. The recombinant molecules can be introduced into the host bacterial cells via transformation, transfection, infection, electroporation, or any other method known in the art.

In a specific embodiment of the invention detailed in the examples herein, plasmids encoding and capable of expressing TMV CP and capable of

**25            5.3.     PRODUCTION AND PURIFICATION OF  
              VIRAL PARTICLES AND OAS-LINKED \*RNA**

Bacterial cells can be grown in a variety of different culture media known in the art, depending on the particular host cell being used. In a specific, preferred embodiment wherein plasmid vectors containing antibiotic resistance genes are employed, the appropriate antibiotics are added depending on the antibiotic resistant genes carried by each of the viral CP and OAS-linked \*DNA vectors. In an embodiment in which the vectors of the invention

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comprise inducible promoters, inducers of the promoter activity, e.g., IPTG for the lac UV5 promoter; are added to the growth media when desired to activate transcription from the vectors.

- 5 Purification of assembled viral particles from bacterial cells can be carried out using a number of different methods. To keep virus-like particles intact during purification, all procedures should avoid shear force and detergents. Standard cleared
- 10 lysate protocols with Triton X-100 should be avoided. Solutions should preferably also contain protease inhibitors such as phenylmethyl sulfonyl fluoride (PMSF), pepstatin, N-tosyl-L-phenylalanine
- 15 chloromethyl ketone (TPCK), and ethylene diamine tetraacetic acid (EDTA) or other divalent cation chelators that would function to inhibit proteases. In particular embodiments, one of the following procedures may be used: (a) osmotic shock in the presence of sucrose (preferred); (b) sucrose pad
- 20 centrifugation of cleared cell lysate; or (c) immunoaffinity with anti-CP antibody. In another embodiment, a freeze-thaw procedure may be useful to lyse the bacterial cells.

- By way of example but not limitation, to
- 25 purify assembled pseudovirus particles, the bacterial cells can be pelleted by centrifugation and then resuspended in a buffered solution such as TE buffer (10 mM Tris-HCl (pH 7.5), 1 mM EDTA) containing lysozyme. After cell lysis, a second centrifugation
- 30 step is carried out to remove cell debris. Pseudovirus particles may be further purified from cleared cell lysate using a number of different methods. For instance, an immunoaffinity column using anti-CP antibodies may be used to further enrich for
- 35 virus particles. Alternatively, the cell lysate may

be centrifuged through a sucrose cushion or gradient and the purified virus particles may be resuspended in TE buffer (10 mM TrisHCl (pH 7.5), 1 mM EDTA). In a preferred aspect, the osmotic shock procedure  
5 described in Section 6.1.2 is used.

To purify the chimeric OAS-linked \*RNA from pseudovirus particles, any procedure known to a skilled artisan may be employed. Preferably, an SDS/phenol extraction is done to remove proteins,  
10 followed by an ethanol precipitation of the RNA.

To confirm that the RNA sequences of interest have been properly expressed and encapsidated within viral CPs according to the present invention, various assays known in the art may be performed,  
15 including but not limited to Northern (dot) blots or PCR (polymerase chain reaction) using reverse transcriptase, using as probe/primers radioactively labeled DNA fragments representative of OAS-linked DNA sequences, or the foreign \*DNA.

20 In some cases, it may be desirable to cleave the OAS-linked sequences from the chimeric RNA prior to use. For example, an OAS sequence located at the 5' end of the RNA could block legitimate translation initiation on \*RNA or initiate translation from an AUG  
25 sequence within the OAS, resulting in a fusion protein which may also not include the desired protein sequences due to initiation not occurring in the proper reading frame. Even if OAS translational initiation sequences can be removed without affecting  
30 packaging function, the presence of the OAS may impede the efficiency of translation of the RNA into the protein of interest. Targeted cleavage, resulting in removal of the OAS sequences, may be accomplished, e.g., by use of RNase H as follows: the OAS-linked  
35 \*RNA is incubated with a DNA oligomer that binds by

virtue of sequence complementarity to the region of the RNA molecule located between the OAS sequence and the RNA sequence of interest. The short region of duplex DNA/RNA that results serves as a substrate for cleavage by the DNA/RNA specific nuclease RNase H. Alternatively, recombinant RNA molecules may be designed to self-cleave by incorporation of a ribozyme domain and recognition sequence for cleavage between the OAS sequence and the DNA sequence of interest. In a different specific embodiment, a sequence-specific cleavage site may be incorporated into the OAS-linked \*RNA in between the OAS and the \*RNA, or a eukaryotic ribosome landing site (IRES) may be inserted between the OAS and \*DNA.

15

#### 5.4. USES OF ENCAPSIDATED RNA

The single stranded RNA packaging system is a convenient and efficient method for production and storage of large amounts of a particular RNA of interest. The system is RNA-sequence and -length independent, and encapsidation of RNA molecules into virus-like particles protects the RNA molecules from substantial degradation by RNases. The pseudovirus particles are easily purified from bacterial cell lysates and, once purified, may be stored for indefinite periods of time in buffer at room temperature. At the desired time, the RNA can be extracted from pseudovirus particles for use.

The uses of the single stranded RNA packaging system are broad and involve almost any of the techniques used in molecular biology that involve manipulation of RNA. For example, in an embodiment where the \*RNA of the OAS-linked \*RNA is a mRNA encoding a protein of interest, the recombinant RNA molecules packaged according to the present invention

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may be released from particles and used in in vitro translation reactions to produce the encoded protein of interest. Alternatively, the packaged recombinant RNA can comprise a sense or antisense RNA molecule that may subsequently be introduced into cells for any of a number of uses, such as to study the possible functions of the encoded protein, for therapeutic effect, etc. The packaging system of the present invention may also be used, by addition of radiolabelled nucleotides to growth media, to synthesize radioactively labeled RNA fragments corresponding to the OAS-linked \*DNA that may be used as hybridization probes, e.g., for northern or Southern blot analysis.

In an alternative embodiment, the pseudovirus particles produced according to the present invention can be introduced directly into animal, plant, fungal, or prokaryotic cells as a means of directly delivering the RNA encapsidated therein, to achieve uncoating and concomitant translation within the cells. Introduction into the cell can be accomplished by any methods known in the art, e.g., by electroporation, or PLO/PEG (poly-L-ornithine/polyethylene glycol) inoculation into spheroplasts or protoplasts in plants. In another embodiment, CP derivatives which contain a surface ligand can effect delivery into animal cells via receptor-mediated endocytosis.

#### 30 5.5. KITS

Kits containing one or more components for carrying out the present invention are also provided. Such a kit comprises in one or more containers component(s) for carrying out the invention. For example, such a kit can comprise container(s) having



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plasmid or other vectors for production of CP and production of OAS-linked \*RNA, as described in Sections 5.1 and 5.2, bacteria containing one or more of the foregoing vectors or chromosomally containing a CP gene or OAS-linked \*DNA. In a specific embodiment, a kit comprises in a first container a bacterium with either a chromosomally integrated or episomal CP gene under the control of preferably an inducible promoter; and in a second container a plasmid vector functional in the bacterial host that is capable of directing the transcription of OAS-linked \*DNA into OAS-linked \*RNA. In a preferred aspect, the second container contains a plasmid vector comprising the following operatively linked components: a promoter, polylinker, OAS, transcriptional termination signal. In one embodiment, the foregoing components are present in 5' to 3' order in which they are listed above. In another embodiment, the plasmid vector comprises in 5' to 3' order: a promoter, OAS, polylinker, transcriptional termination signal.

## 6. EXAMPLES

### 6.1. MATERIALS AND METHODS

#### 6.1.1. PLASMIDS AND BACTERIAL STRAINS

The host bacterial strain was *E. coli* BL21 (DE3). *E. coli* BL21 is F<sup>+</sup>, ompT, r<sub>g</sub><sup>-</sup>, mβ<sup>-</sup>. DE3 is a λ derivative which carries a DNA fragment containing the lacI gene, the lac UV5 promoter, the beginning of the lacZ gene, and the gene for T7 RNA polymerase. The bacterial strain was originally provided by the Brookhaven National Research Laboratory and is a λ lysogen in which the structural gene for bacteriophage T7 RNA polymerase is expressed from the lac UV5 promoter when the inducer IPTG is added to the medium (Studier et al., 1990, Meth. Enzymol. 185:62-89). The

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lysogenic host strain also contains the plasmid vector pACYC184 (Pouwels et al., 1985, in *Cloning Vectors*, Elsevier Science Publishers, Amsterdam, p. I-A-iv-9) which is a p15A replicon and, as pLySE (Studier et al., 1990, *Meth. Enzymol.* 185:60-89), carries the T7 lysozyme structural gene expressed from the tet promoter. The lysozyme functions to bind to the T7 RNA polymerase and inhibit the low level of constitutive transcription which occurs from the lac UV5 promoter even in the absence of the IPTG inducer. The lysozyme also makes the cells more fragile since it degrades the peptidoglycan wall of the *E. coli* cells.

As part of the expression system, a source of TMV CP must be supplied. Various TMV CP structural genes (including derivatives thereof) were cloned into plasmid vector pET3a (Fig. 6). pET3a contains a T7 promoter from phage T7 gene 10 as well as the translational initiation signal (including the prokaryotic Shine-Dalgarno site) from gene 10 of this phage. The pET3 plasmid series also contains a T7 RNA polymerase terminator and is a ColE1 replicon, which is compatible with the pACYC184/p15A replicon described above. Between the T7 promoter and terminator are *NdeI* and *NheI* sites for the insertion and cloning of sequences to be transcribed by the T7 RNA polymerase.

An expression plasmid for production of native TMV coat protein using the native codons of the eukaryotic virus was made by digesting a full-length clone of TMV in the plasmid pTMV210 (a gift from W.O. Dawson, Lake Alfred University of Florida) with *NsiI*, followed by treatment with T4 DNA polymerase to "chew back" 3' protruding termini, and subsequent digestion with *DraI*. The resulting fragment was inserted into

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*Sma*I-digested pGEM3Z (Promega, Madison, WI), and the resulting KpnI-BamHI CP-containing fragment was inserted (after trimming of the KpnI site to make blunt ends) into pET3a. The resulting plasmid, pET301, produces the native TMV CP (encoding the N-terminal Met-Ser of the U1 strain of TMV) from a T7 promoter when IPTG is added to induce high level expression of the T7 RNA polymerase gene (from DE3). Another plasmid (kindly provided by Joel Haynes) produces the native TMV CP of the U1 strain (Haynes et al., 1986, *Biotechnology* 4:637-641). This latter plasmid contains an entirely synthetic TMV CP gene insofar as the codons have been optimized for *E. coli* expression. Once again, an *Eco*RI-BamHI cassette of this gene was trimmed down and cloned into pET3a to produce pET302.

Plasmids were also constructed that coded for coat proteins in which the serine at the second amino acid position was replaced by alanine, aspartic acid or proline. The resulting plasmids were designated pETAla301, pETAsp301 and pETPro301, respectively. The site directed mutagenesis of the 5' end of the gene encoding TMV coat protein was done by PCR reactions using the following primers:

1. for changing Ser(2) to Ala; 5' primer 5'-GCCATGGCTTACAGTATCACTACT-3' (SEQ ID NO:1); 3' primer 5'-GGTCGACCTCAAGTTGCAGGACCA-3' (SEQ ID NO:2).
2. for changing Ser(2) to Asp; 5' primer 5'-GCCATGGACTACAGTATCACTACT-3' (SEQ ID NO:3); 3' primer was that of SEQ ID NO:2.
3. for changing Ser(2) to Pro; 5' primer 5'-GGCATGCCGTACAGTATCACTACT-3' (SEQ ID NO:4); 3' primer was that of SEQ ID NO:2.

Two strategies were used to create substrate transcripts containing the TMV origin-of-assembly (OAS) sequence for encapsidation by TMV coat protein *in situ*. The first system used the chloramphenicol acetyl transferase (CAT; Cm') gene in the plasmid pLysE, which also harbors the T7 lysozyme gene. The unique EcoRI site in the Cm' gene was cleaved and an EcoRI fragment from plasmid pJII102 (Gallie et al., 1987, Science 236:1122-24) containing the 3' half of the chloramphenicol acetyl transferase (Cm') gene and the TMV assembly origin (OAS) (an -440 nucleotide sequence, from bases 5118-5550 of the viral RNA sequence; Zimmermann, 1983, EMBO J. 2:1901-07; Fig. 10, SEQ ID NO:5) was inserted into the opened EcoRI site of pLysE to generate pLys102 (Fig. 7). Thus, low level constitutive transcription from the *E. coli* promoter of the Cm' gene, used as a selectable marker for the plasmid pLysE, created a functional chloramphenicol acetyl transferase mRNA followed by the TMV OAS, in turn followed by the repeated remnant 3' terminal fragment of the original chloramphenicol acetyl transferase selectable marker gene. Transcripts were thus expected to be 1.7 kb in length with an internal OAS.

A second approach to create OAS<sup>+</sup> RNA for encapsidation with TMV coat protein *in situ* used a T7 promoter-poly (CAA) leader-GUS gene cassette derived from the plasmid pJII1032 (a pUC19 derivative which contain *E. coli*  $\beta$ -glucuronidase (GUS); see Wilson, T.M.A., et al., 1990, in *Post-Transcriptional Control of Gene Expression*, McCarthy and Tuite (eds.), NATO-ASI Series, Vol. H49, pp. 261-75), to which was attached a Sall-HindIII cassette containing the TMV OAS (an -440 nucleotide sequence, from bases 5118-5550 of the viral RNA sequence; Zimmermann, 1983, EMBO J.

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2:1901-07; Fig. 10; SEQ ID NO:5) and a T7 terminator at the 3' side of the GUS gene. The whole *EcoRI*-*HindIII* fragment was then modified and cloned into a unique *HindIII* site in the plasmid pLysE to create  
5 pLys1032 (Fig. 8). Transcripts derived from the T7 promoter are expected to be approximately 2.5 kb in length and contain a 3' terminal TMV origin-of-assembly sequence.

10 6.1.2. PURIFICATION OF PSEUDOVIRAL PARTICLES FROM *E. COLI*

Bacteria were grown in 100 ml of media at 37°C for 4 hours. Cells were induced by 0.4 mM IPTG for 6 hours at room temperature. Bacteria were  
15 harvested by centrifugation at 8K rpm at 4°C for 15 minutes. The bacterial cell pellet was resuspended in 1 ml of TE (10 mM Tris-HCl (pH 7.5), 1 mM EDTA) and then lysozyme solution was added to 1 mg/ml. This suspension was incubated at RT for 5-10 minutes  
20 followed by addition of four volumes of TE (pH 7.5). DNase I was added and the suspension was incubated until it was no longer viscous. The suspension was centrifuged at 10 K rpm, at 4°C for 20 minutes to remove cell debris and the supernatant was loaded onto  
25 a sucrose gradient (10-40%) and spun in a Beckman 45 Ti rotor at 40K rpm, at 4°C for 4 hours. The pellet from the bottom of the sucrose gradient was dissolved in 1 ml of TE (pH 7.5) and this sample was used for electron microscopy.

30

6.1.3. EXTRACTION OF RNA AND NORTHERN DOT BLOTS

A <sup>32</sup>P-labelled DNA fragment containing the OAS sequence was used as a probe in northern dot blot  
35 hybridizations. The probe was derived from pJII102

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- (Gallie et al., 1987, Science 236:1122-24), a portion of which is shown in Figure 9. pJII102 was digested with BamHI, run on a gel, and a small, approximately 440 bp fragment containing OAS was eluted. The
- 5 fragment was nick-translated with  $^{32}\text{P}$ -dATP using *E. coli* DNA polymerase I and DNase I (see Sambrook et al., 1989, Molecular Cloning, A Laboratory Manual, 2nd Ed., Cold Spring Harbor Laboratory, N.Y.) The labeled probe was added to the hybridization solution.
- 10 For the dot blot hybridization, RNA was extracted from purified viral particles by phenol extraction. The RNA was precipitated by the addition of ethanol and an appropriate salt such as sodium acetate. The RNA sample was dotted onto
- 15 nitrocellulose paper, which was then dried at room temperature. After denaturation and neutralization, the filter was baked at 80°C for 2 h, prehybridized at 42°C for 2 h, and then hybridized at 42°C for 2 h with the nick-translated probe, washed, and exposed to
- 20 film.

6.1.4. POLYMERASE CHAIN REACTION  
(PCR) AMPLIFICATION OF RNA  
ISOLATED FROM PSEUDOVIRUS PARTICLES

- RNA samples were extracted from purified
- 25 TMV-like particles as described in Section 6.1.3. The RNA samples were used as substrates for reverse transcriptase, and the resulting cDNAs were used in PCR reactions with the following pair of primers:
- 3' primer of TMV OAS (5' to 3'),
- 30 5' CCG GTT CGA GAT CGA 3' (SEQ ID NO:7); and 5' primer of TMV OAS (5' to 3'), 5' GTT GGT CGT CAC GG 3' (SEQ ID NO:8).

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#### 6.1.5. SDS PAGE ANALYSIS OF E. COLI PRODUCED TMV COAT PROTEIN

The TMV CP used in these particular experiments was either the *E. coli* optimized codon version of the U1 strain sequence (pET302) or native TMV U1 coat protein sequence (pET301) or the native TMV coat protein sequence PCR cloned from pTMV210 but in which the second amino acid (serine) was altered to an alanine (pETAla301). IPTG was added to the 10 ml culture when the cell density reached approximately 0.6. In the cases shown, the induction time with IPTG was 2 hours. A volume equivalent to 93  $\mu$ l of original *E. coli* culture was loaded on a standard discontinuous Laemmli gel system (Laemmli, U.K., 1970, Nature 227:680) in which the resolving gel was 15% polyacrylamide and subjected to sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS PAGE). To visualize the proteins, the gel was stained with Coomassie blue. The identity of the protein bands was confirmed not only by co-migration with legitimate TMV CP (added to the marker lane 2) but also by western blotting with polyclonal antiserum raised in rabbits against native U1 TMV CP.

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#### 6.2. RESULTS

##### 6.2.1. EXPRESSION OF TMV VIRAL COAT PROTEINS AND PACKAGING OF RECOMBINANT RNA MOLECULES IN E. COLI

The results of SDS PAGE analysis of cleared cell lysates are shown in Figure 1. The TMV coat protein used in these particular experiments was either the *E. coli* optimized codon version of the U1 strain sequence (pET302) or the native TMV U1 coat protein sequence (pET301) or the native TMV coat protein sequence PCR cloned from pTMV210 but in which the second amino acid (serine) is altered to an

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alanine. Staining with Coomassie blue clearly shows that pET301 and pETAla301 induced synthesis of 2-3  $\mu$ g TMV coat protein per 93  $\mu$ l of original cell culture. pET302 induced about 2 or 3 times more CP than either  
5 of the above as judged by Coomassie blue staining. The identity of these protein bands was confirmed not only by co-migration with legitimate TMV coat protein (added to the marker lane 2) but also by western  
10 blotting with polyclonal antiserum raised in rabbits against TMV coat protein (Fig. 2). The low level of signal on the Western blot in lane 2 is attributed to the fact that, even without IPTG, a low level  
background constitutive expression of the T7 RNA  
polymerase takes place despite the expression of T7  
15 lysozyme which inhibits the activity of the enzyme.

For purposes of comparison to the *in vivo* assembly of particles, *in vitro* assembly was also attempted. All the CPs were purified extensively from induced cultures of *E. coli* in the absence of OAS+  
20 messenger RNAs and following the procedure of Durham (1972, J. Mol. Biol. 67:289-305), were incubated with phenol extracted purified TMV RNA. Virus-purified CP was also used as a positive control. The positive control assembled and increased turbidity ( $A_{310}$  nm)  
25 measurably, whereas none of the *E. coli* generated CPs functioned to assemble with TMV RNA *in vitro*.

Electron micrographs of *E. coli* in which each CP construct had been expressed in the absence of an OAS+ RNA revealed some assembly of virus-like  
30 structures which, upon close examination, were seen to be predominantly stacked disk rods (usually viewed as products of proteolysis) which presumably contain no RNA. This observation of non-helical assembly has been confirmed by the absence of immunogold labelling

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with monoclonal antibodies specific for the native helical TMV structure.

With regard to *in vivo* particle assembly in *E. coli*, in both the presence or absence of OAS+ mRNA, the proline version of the TMV coat protein, synthesized from pETPro301 clearly formed stacked disk rods as visualized in the electron microscope. We have yet to determine whether or not these rods contain RNA but their clear, striated appearance suggests a protein-only structure. Cleared cell lysates were viewed in the electron microscope and the resulting pictures are shown in Figure 3 A-D. The native form of TMV CP as generated from plasmid pET301, when expressed without an OAS+ RNA, gave a completely clean background and no apparent assembly into virus-like rods. For pETAla301, pET301, and pET302, when coat protein from any one of these plasmids was expressed in the presence of an OAS+ mRNA, for either CAT or GUS, electron microscopy revealed a substantial increase in the number of short rods, as demonstrated in Figure 3 for pETAla301. The lengths of some preparations were more homogeneous than others and complied with the predicted lengths of the rods (approximately 75 nm for CAT-OAS) resulting from transcription of the particular gene.

RNA was extracted from these purified particles following sucrose density gradient, cesium sulfate isopycnic gradient or sucrose pad centrifugation, and a northern dot blot was done with a TMV OAS specific probe. In the case of induction where CAT and/or GUS RNA containing the TMV origin-of-assembly sequence together with pETAla301 coat protein was expressed, the resulting rods produced RNA which gave a strong and clear signal with an OAS specific probe (Fig. 4). PCR amplification of RNA samples

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derived from pseudovirus particles by use of 5' and 3' primers from the TMV OAS sequences also allowed detection of CAT-OAS and GUS-OAS in such case (Fig. 5).

5                   Thus, pETAla301-encoded TMV CPs assembled and packaged CAT-OAS RNA or CAA-GUS-OAS RNA into TMV-like pseudovirus particles *in vivo* in *Escherichia coli*.

10                   7. DEPOSIT OF MICROORGANISMS

Bacteria strain *E. coli* BL21 (DE3) containing plasmid pETAla301 (an expression plasmid encoding a TMV coat protein with Ser<sub>2</sub> replaced by Ala<sub>2</sub>), and plasmid pLys102 (a plasmid providing  
15                   packageable CAT-OAS transcripts) was deposited on September 10, 1992 with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852, under the provisions of the Budapest Treaty on the International Recognition of the Deposit of  
20                   Microorganisms for the Purposes of Patent Procedures, and assigned Accession No. 69070.

The present invention is not to be limited in scope by the microorganism deposited or the specific embodiments described herein. Indeed,  
25                   various modifications of the invention in addition to those described herein will become apparent to those skilled in the art from the foregoing description and accompanying figures. Such modifications are intended to fall within the scope of the appended claims.

30                   Various publications are cited herein, the disclosures of which are incorporated by reference in their entireties.

- 40 -

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT: Wilson, Thomas M.A. et al.
- (ii) TITLE OF INVENTION: RNA Packaging System
- (iii) NUMBER OF SEQUENCES: 8
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Pennie & Edmonds
  - (B) STREET: 1155 Avenue of the Americas
  - (C) CITY: New York
  - (D) STATE: New York
  - (E) COUNTRY: U.S.A.
  - (F) ZIP: 10036-2711
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER:
  - (B) FILING DATE: On Even Date Herewith
  - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Coruzzi, Laura A.
  - (B) REGISTRATION NUMBER: 30,742
  - (C) REFERENCE/DOCKET NUMBER: 7108-006
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: 212 790-9090
  - (B) TELEFAX: 212 869-9741
  - (C) TELEX: 66141 PENNIE

## (2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 24 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: DNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

GCCATGGCTT ACAGTATCAC TACT

24

## (2) INFORMATION FOR SEQ ID NO:2:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 24 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: DNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

- 41 -

GGTCGACCTC AAGTTGCAGG ACCA

24

## (2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 24 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: DNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

GCCATGGACT ACACTATCAC TACT

24

## (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 24 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: DNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

GGCATGCCGT ACACTATCAC TACT

24

## (2) INFORMATION FOR SEQ ID NO:5:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 433 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: RNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GGUCGUCACG GGCGAGUGGA ACUUGCCUGA CAAUUGCAGA GGAGGUGUGA GCGUGUGUCU	60
GGUGGACAAA AGGAUGGAAA GAGCCGACGA GGCCACUCUC GGAUCUUAU ACACAGCAGC	120
UGCAAAGAAA AGAUUUCAGU UCAAGGUGGU UCCCAAUUU GCUAUAACCA CCCAGGACGC	180
GAUGAAAAAC GUCUGGCAAG UUUUAGUUA UAUUAGAAU GUGAAGUGU CAGCCGGUUU	240
CUGUCCGCUU UCUCUGGAGU UUGUGUCGGU GUGUAUUGU UAUAGAAUA AUUAUAAAUU	300
AGGUUUGAGA GAGAAGAUUA CAAACGUGAG AGACGGAGGG CCCAUGGAAC UUACAGAAGA	360
AGUCGUUGAU GAGUUAUGG AAGAUGUCCC UAUGUCGAUC AGGCUUGCAA AGUUUCGAUC	420
UCCAACCGGA AAA	433

## (2) INFORMATION FOR SEQ ID NO:6:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 75 nucleotides
  - (B) TYPE: nucleic acid

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- (C) STRANDEDNESS: single
- (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: RNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

UGAGAGACGG AGGGCCCAUG GAACUUACAG AAGAAGUCGU UGAUGAGUUC AUGGAAGAUG 60  
UCCCUAUGUC GAUCA 75

(2) INFORMATION FOR SEQ ID NO:7:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 15 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: DNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

CCGGTTCGAG ATCGA 15

(2) INFORMATION FOR SEQ ID NO:8:

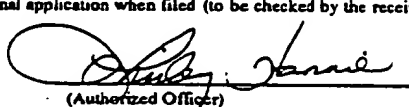
- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 14 nucleotides
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: DNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

GTTGGTCGTC ACGG 14

International Application No: PCT/

MICROORGANISMS	
Optional Sheet in connection with the microorganism referred to on page 39, lines 10-21 of the description *	
<b>A. IDENTIFICATION OF DEPOSIT *</b> Further deposits are identified on an additional sheet *	
Name of depositary institution * American Type Culture Collection	
Address of depositary institution (including postal code and country) * 12301 Parklawn Drive Rockville, MD 20852 US	
Date of deposit * <u>September 10, 1992</u> Accession Number * <u>69070</u>	
<b>B. ADDITIONAL INDICATIONS</b> * (leave blank if not applicable). This information is contained on a separate attached sheet	
<b>C. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE *</b> * of the indications are not all designated States	
<b>D. SEPARATE FURNISHING OF INDICATIONS *</b> (leave blank if not applicable) The indications listed below will be submitted to the International Bureau later * (Specify the general nature of the indications e.g., "Accession Number of Deposit")	
<b>E.</b> <input checked="" type="checkbox"/> This sheet was received with the International application when filed (to be checked by the receiving Office) <div style="text-align: right;"> (Authorized Officer)</div> <input type="checkbox"/> The date of receipt (from the applicant) by the International Bureau * was _____ (Authorized Officer)	

WHAT IS CLAIMED IS:

1. A method of producing and encapsidating a recombinant RNA molecule in a plant pseudovirus particle comprising culturing a bacterium containing  
5 (a) a first recombinant nucleic acid encoding a plant virus coat protein or functional derivative thereof, and (b) a second recombinant nucleic acid that can be transcribed to produce an RNA molecule comprising the  
10 plant virus origin-of-assembly sequence operatively linked to an RNA sequence of interest, with the proviso that the RNA sequence of interest is not the plant virus RNA; whereby the plant virus coat protein is expressed and the second recombinant nucleic acid  
15 is transcribed in the bacterium, and the coat protein assembles into a particle encapsidating the RNA molecule; in which the plant virus has a rod-shaped helical particle and a single-stranded RNA genome.
- 20 2. The method according to claim 1 in which the plant virus is selected from the group consisting of a tobamovirus, a potexvirus, a tobnavirus, a hordeivirus, a potyvirus, and a furovirus.
- 25 3. The method according to claim 2 in which the plant virus is a tobamovirus.
4. The method according to claim 3 in  
30 which the plant virus is tobacco mosaic virus.
5. The method according to claim 1 in which the first recombinant nucleic acid comprises a promoter sequence which controls the expression of the  
35 plant virus coat protein in the bacterium; and the

second recombinant nucleic acid comprises a promoter sequence which controls the expression of the RNA in the bacterium.

5                   6. The method according to claim 1 in which the first recombinant nucleic acid is a plasmid vector comprising the following operatively linked components:

- 10                   (a) a promoter which controls the expression of the coat protein or functional derivative thereof;
- (b) a translation initiation signal;
- (c) a DNA sequence encoding the coat protein or functional derivative thereof; and
- 15                   (d) a transcription termination signal.

                  7. The method according to claim 4 in which the first recombinant nucleic acid is a plasmid vector comprising the following operatively linked components:

- 25                   (a) a promoter which controls the expression of the coat protein or functional derivative thereof;
- (b) a translation initiation signal;
- (c) a DNA sequence encoding the coat protein or functional derivative thereof; and
- 30                   (d) a transcription termination signal.

                  8. The method according to claim 1 in which the second recombinant nucleic acid is a plasmid

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vector comprising the following operatively linked components:

- (a) a promoter;
- (b) a DNA sequence that can be transcribed under the control of the promoter in the bacterium to produce the RNA molecule comprising the origin of assembly sequence operatively linked to the RNA sequence of interest; and
- (c) a transcription termination signal.

9. The method according to claim 6 in which the second recombinant nucleic acid is a plasmid vector comprising the following operatively linked components:

- (a) a promoter;
- (b) a DNA sequence that can be transcribed under the control of the promoter in the bacterium to produce the RNA molecule comprising the origin of assembly sequence operatively linked to the RNA sequence of interest; and
- (c) a transcription termination signal.

10. The method according to claim 9 in which the first recombinant nucleic acid and the second recombinant nucleic acid each further comprises a selectable marker.

11. The method according to claim 10 in which the selectable marker is an antibiotic resistance gene.

5 12. The method according to claim 10 in which the bacterium further contains a recombinant DNA sequence which is expressed to produce lysozyme upon culturing the bacterium.

10 13. The method according to claim 4 in which the first recombinant nucleic acid comprises the native coat protein gene of a tobacco mosaic virus.

15 14. The method according to claim 4 in which the first recombinant nucleic acid comprises a sequence encoding a native coat protein of a tobacco mosaic virus, which sequence has been optimized for translation in *Escherichia coli*.

20 15. The method according to claim 4 in which the first recombinant nucleic acid comprises a sequence encoding the coat protein of the U1 strain of tobacco mosaic virus in which the second amino acid from the amino termini has been changed to an alanine.

25 16. The method according to claim 4 in which the origin of assembly sequence comprises the sequence depicted in Figure 11 (SEQ ID NO:6).

30 17. The method according to claim 15 in which the origin of assembly sequence comprises the sequence depicted in Figure 11 (SEQ ID NO:6).

18. The method of claim 4 in which the origin of assembly sequence comprises the sequence depicted in Figure 10 (SEQ ID NO:5).

5 19. A method of producing and encapsidating a recombinant RNA molecule in a plant pseudovirus particle comprising culturing a bacterium containing (a) a first plasmid vector comprising the following operatively linked components: (i) a first promoter  
10 which controls the expression of a first DNA sequence encoding a tobacco mosaic virus coat protein or functional derivative thereof, (ii) a translation initiation signal, (iii) the first DNA sequence encoding the coat protein or functional derivative  
15 thereof, and (iv) a first transcription termination signal; and (b) a second plasmid vector comprising the following operatively linked components: (i) a second promoter, (ii) a second DNA sequence that can be transcribed under the control of the second promoter  
20 in the bacterium to produce an RNA molecule comprising a tobacco mosaic virus origin of assembly sequence operatively linked to an RNA sequence of interest, with the proviso that the RNA sequence is not a tobacco mosaic virus RNA, and (iii) a second  
25 transcription termination signal; whereby the coat protein or derivative thereof is expressed and the second DNA sequence is transcribed in the bacterium, and the coat protein or derivative thereof assembles into a particle encapsidating the RNA molecule.  
30

20. The method according to claim 19 in which the first DNA sequence encodes the coat protein of the U1 strain of tobacco mosaic virus in which the second amino acid from the amino terminus has been  
35 changed to an alanine, and the origin of assembly

sequence comprises the sequence depicted in Figure 11  
(SEQ ID NO:6).

21. The method according to claim 20 in  
5 which the bacterium is an *Escherichia coli*.

22. The method according to claim 19 in  
which the first promoter and the second promoter are  
inducible promoters.

10

23. The method according to claim 19 in  
which the first promoter and the second promoter are  
T7 RNA promoters, and the bacterium further contains a  
third DNA sequence comprising a T7 RNA polymerase  
15 coding sequence and an inducible promoter controlling  
the expression of the T7 RNA polymerase.

20

24. The method according to claim 23 in  
which the inducible promoter is a lac UV5 promoter.

25. The method according to claim 24 in  
which the third DNA sequence is chromosomally  
integrated.

25

26. The method according to claim 1 in  
which the RNA sequence of interest is a messenger RNA  
sequence encoding a protein of interest.

27. The method according to claim 19 in  
30 which the RNA sequence of interest is a messenger RNA  
sequence encoding a protein of interest.

35

28. A bacterium containing:  
(a) a first recombinant nucleic acid  
encoding a plant virus coat

50

protein or functional derivative thereof; and

- (b) a second recombinant nucleic acid that can be transcribed in a suitable bacterium to produce an RNA molecule comprising the plant virus origin of assembly sequence operatively linked to an RNA sequence of interest, with the proviso that the RNA sequence of interest is not the plant virus RNA,

in which the plant virus has a rod-shaped helical particle and a single-stranded RNA genome.

29. The bacterium of claim 28 in which the plant virus is selected from the group consisting of a tobamovirus, a potexvirus, a tobnavirus, a hordeivirus, a potyvirus, and a furovirus.

30. The bacterium of claim 29 in which the plant virus is tobacco mosaic virus.

31. The bacterium of claim 28 in which the first recombinant nucleic acid comprises a promoter sequence which controls the expression of the plant virus coat protein in the bacterium; and the second recombinant nucleic acid comprises a promoter sequence which controls the expression of the RNA molecule in the bacterium.

32. The bacterium of claim 28 in which the first recombinant nucleic acid encodes the coat protein of the U1 strain of tobacco mosaic virus in which the second amino acid from the amino termini has

been changed to an alanine, and in which the origin of assembly sequence comprises the sequence depicted in Figure 11 (SEQ ID NO:6).

- 5                   33. A recombinant bacterium containing:
- (a) a first plasmid vector comprising
- the following operatively linked
- components: (i) a first promoter
- 10                   which controls the expression of a
- first DNA sequence encoding a
- tobacco mosaic virus coat protein
- or functional derivative thereof;
- (ii) a translation initiation
- signal; (iii) the first DNA
- 15                   sequence encoding the coat protein
- or derivative thereof; and (iv) a
- first transcription termination
- signal; and
- (b) a second plasmid vector comprising
- 20                   the following operatively linked
- components: (i) a second promoter;
- (ii) a second DNA sequence that
- can be transcribed under the
- control of the second promoter in
- 25                   the bacterium to produce an RNA
- molecule comprising a tobacco
- mosaic virus origin of assembly
- sequence operatively linked to an
- RNA sequence of interest, with the
- 30                   proviso that the RNA sequence is
- not a tobacco mosaic virus RNA;
- and (iii) a second transcription
- termination signal;

                  whereby the coat protein or derivative thereof is

35                   expressed and the second DNA sequence is transcribed

in the bacterium, and the coat protein or derivative thereof assembles into a particle encapsidating the RNA molecule.

5           34. The bacterium of claim 33 in which the first DNA sequence encodes the coat protein of the U1 strain of tobacco mosaic virus in which the second amino acid from the amino terminus has been changed to an alanine, and in which the origin of assembly  
10 sequence comprises the sequence depicted in Figure 11 (SEQ ID NO:6).

          35. The bacterium of claim 34 in which the bacterium is an *Escherichia coli*.

15           36. The bacterium of claim 33 in which the first promoter and the second promoter are T7 RNA promoters, and the bacterium further contains a third DNA sequence comprising a T7 RNA polymerase coding  
20 sequence and an inducible promoter controlling the expression of the T7 RNA polymerase.

          37. An *Escherichia coli* containing plasmids pETAla301 and pLys102 as deposited with the ATCC and  
25 assigned Accession No. 69070.

          38. A kit comprising, in one container:  
          (a) a first recombinant nucleic acid  
              comprising a DNA sequence encoding  
30 a plant virus coat protein or functional derivative thereof and a promoter sequence which controls the expression of the plant virus coat protein in a suitable  
35 bacterium; and

## 53

- 5 (b) a second recombinant nucleic acid that can be transcribed in the suitable bacterium to produce an RNA molecule comprising the plant virus origin of assembly sequence operatively linked to an RNA sequence of interest, with the proviso that the RNA sequence of interest is not the plant virus
- 10 RNA;

in which the plant virus has a rod-shaped helical particle and a single-stranded RNA genome.

39. The kit of claim 38 in which the plant
- 15 virus is selected from the group consisting of a tobamovirus, a potexvirus, a tobnavirus, a hordeivirus, a potyvirus, and a furovirus.

40. The kit of claim 39 in which the plant
- 20 virus is tobacco mosaic virus.

41. The kit of claim 40 in which the first recombinant nucleic acid encodes the coat protein of the U1 strain of tobacco mosaic virus in which the
- 25 second amino acid from the amino termini has been changed to an alanine, and in which the origin of assembly sequence comprises the sequence depicted in Figure 11 (SEQ ID NO:6).

- 30 42. A kit comprising, in one container:
- (a) a first plasmid vector comprising the following operatively linked components: (i) a first promoter which controls the expression of a
- 35 first DNA sequence encoding a



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- 5 tobacco mosaic virus coat protein  
or functional derivative thereof;  
(ii) a translation initiation  
signal; (iii) the first DNA  
sequence encoding the coat protein  
or derivative thereof; and (iv) a  
first transcription termination  
signal; and
- 10 (b) a second plasmid vector comprising  
the following operatively linked  
components: (i) a second promoter;  
(ii) a second DNA sequence that  
can be transcribed under the  
control of the second promoter in  
15 the bacterium to produce an RNA  
molecule comprising a tobacco  
mosaic virus origin of assembly  
sequence operatively linked to an  
RNA sequence of interest, with the  
proviso that the RNA sequence is  
20 not a tobacco mosaic virus RNA;  
and (iii) a second transcription  
termination signal;

whereby the coat protein or derivative thereof is  
25 expressed and the second DNA sequence is transcribed  
in the bacterium, and the coat protein or derivative  
thereof assembles into a particle encapsidating the  
RNA molecule.

- 30 43. The kit of claim 42 in which the first  
DNA sequence encodes the coat protein of the U1 strain  
of tobacco mosaic virus in which the second amino acid  
from the amino terminus has been changed to an  
alanine, and in which the origin of assembly sequence  
35

55

comprises the sequence shown in Figure 11 (SEQ ID NO:6).

44. The kit of claim 42 in which the  
5 bacterium is an *Escherichia coli*.

45. The kit of claim 42 in which the first  
promoter and the second promoter are T7 RNA promoters,  
and the bacterium further contains a third DNA  
10 sequence comprising a T7 RNA polymerase coding  
sequence and an inducible promoter controlling the  
expression of the T7 RNA polymerase.

46. A kit comprising, in one container:  
15 (a) a first plasmid vector comprising  
the following operatively linked  
components: (i) a first promoter  
which controls the expression of a  
first DNA sequence encoding a  
20 tobacco mosaic virus coat protein  
or functional derivative thereof;  
(ii) a translation initiation  
signal; (iii) the first DNA  
sequence encoding the coat protein  
25 or derivative thereof; and (iv) a  
first transcription termination  
signal; and  
(b) a second plasmid vector comprising  
the following operatively linked  
30 components: (i) a second promoter  
which controls the expression of a  
second DNA sequence, (ii) the  
second DNA sequence, that  
comprises a polylinker and a  
35 sequence encoding a tobacco mosaic

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virus origin of assembly sequence,  
and (iii) a second transcription  
termination signal.

5           47. The kit according to claim 46 in which  
the polylinker is 5' to the sequence encoding a  
tobacco mosaic virus origin of assembly sequence.

10           48. The kit according to claim 46 in which  
the polylinker is 3' to the sequence encoding a  
tobacco mosaic virus origin of assembly sequence.

15           49. A kit comprising in one container:  
(a) a first plasmid vector comprising  
the following operatively linked  
components: (i) a first promoter  
which controls the expression of a  
first DNA sequence encoding a  
plant virus coat protein or  
20 functional derivative thereof;  
(ii) a translation initiation  
signal; (iii) the first DNA  
sequence encoding the coat protein  
or derivative thereof; and (iv) a  
25 first transcription termination  
signal; and  
(b) a second plasmid vector comprising  
the following operatively linked  
components: (i) a second promoter  
30 which controls the expression of a  
second DNA sequence, (ii) the  
second DNA sequence, that  
comprises a polylinker and a  
sequence encoding the plant virus  
35 origin of assembly sequence, and

57

(iii) a second transcription termination signal;  
in which the plant virus has a rod-shaped helical particle and a single-stranded RNA genome.

5

50. A kit comprising in one or more containers the bacterium of claim 28.

51. A kit comprising in one or more  
10 containers the bacterium of claim 33.

52. A kit comprising in one or more containers the bacterium of claim 36.

15

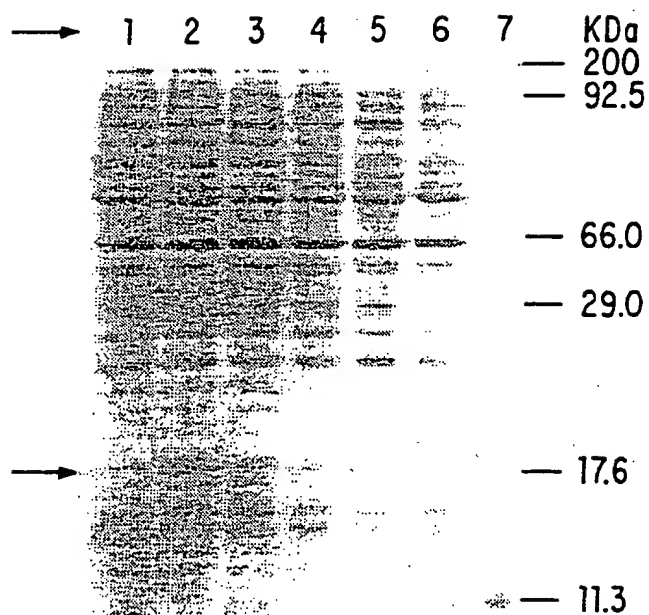
20

25

30

35

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**FIG. 1**

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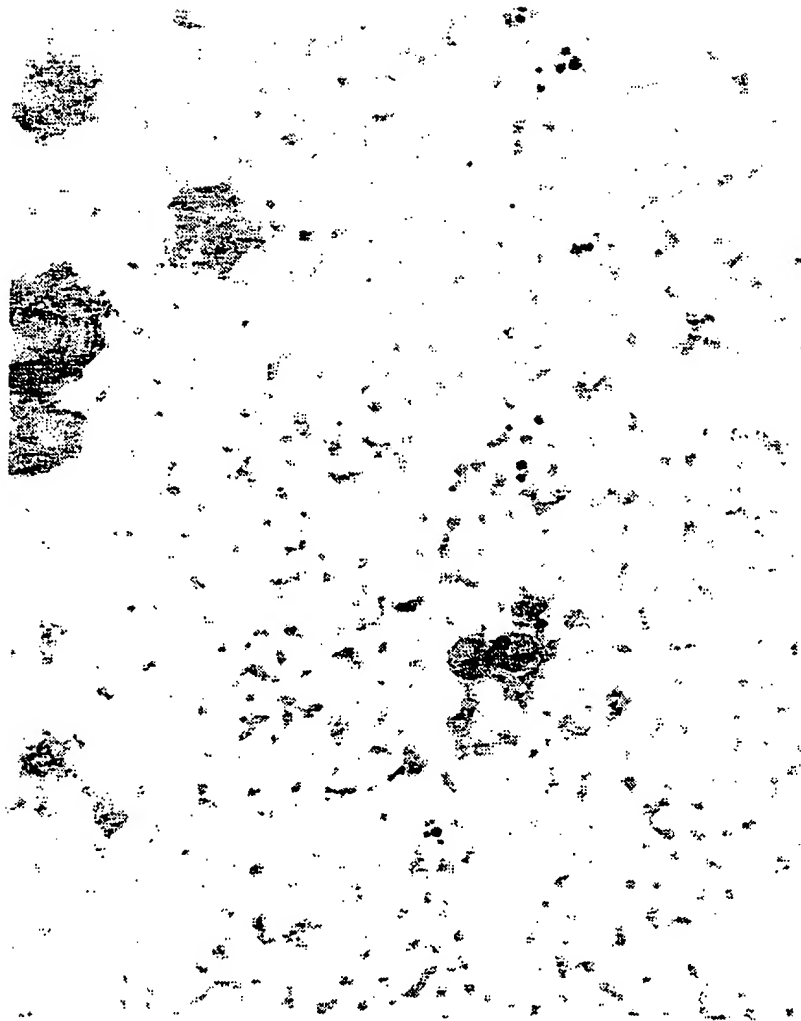
1 2 3 4

← TMV CP

**FIG. 2**

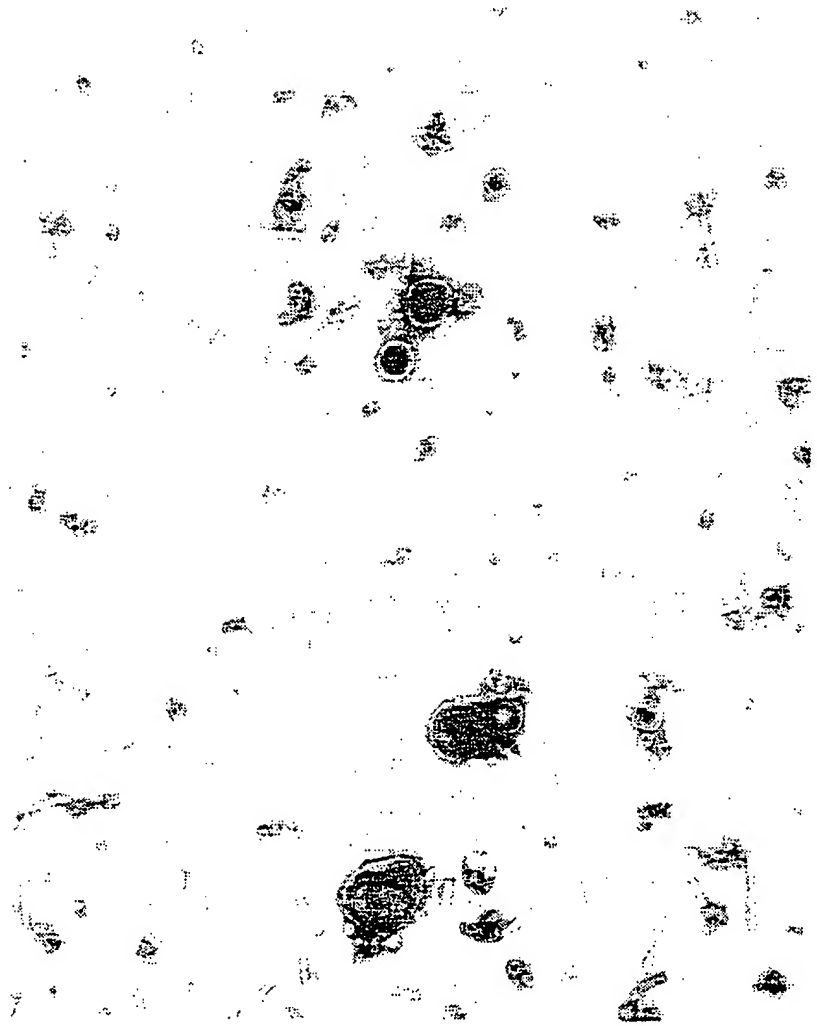
SUBSTITUTE SHEET (RULE 26)

3/12



**FIG. 3A**

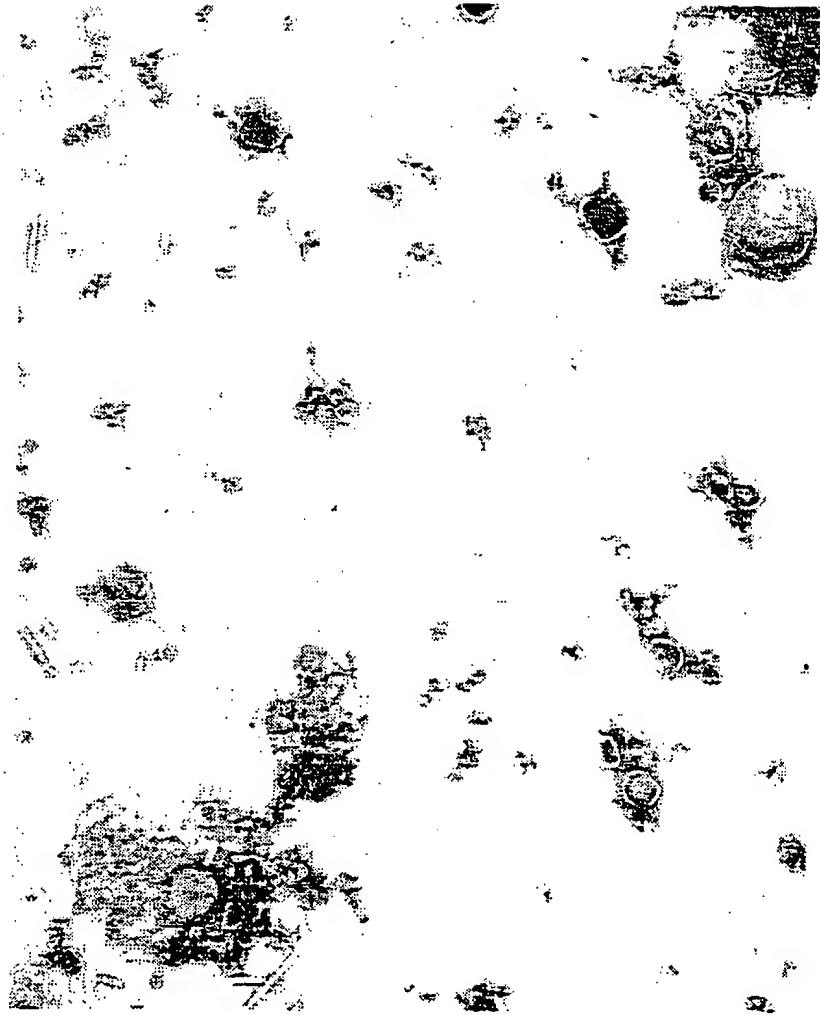
4/12



**FIG. 3B**



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**FIG. 3C**

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**FIG. 3D**

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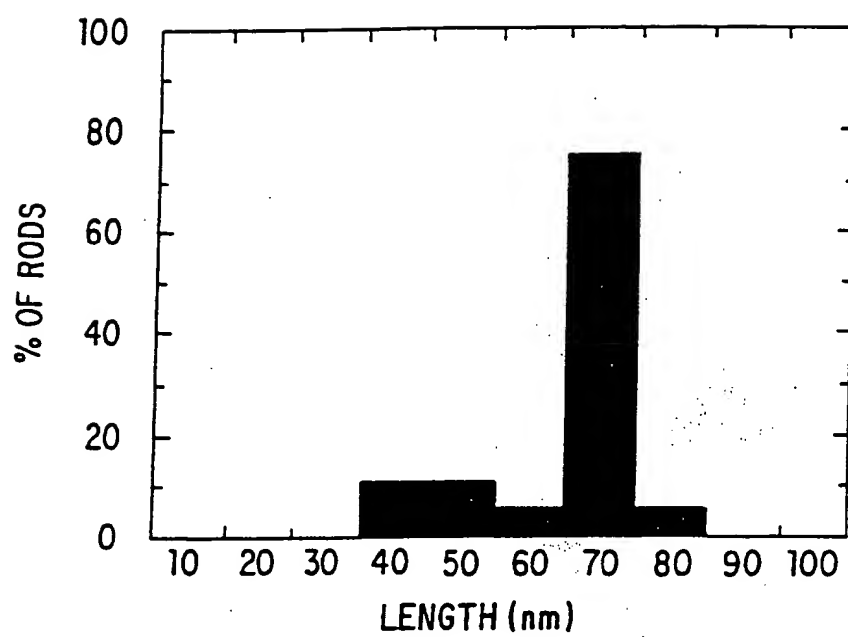
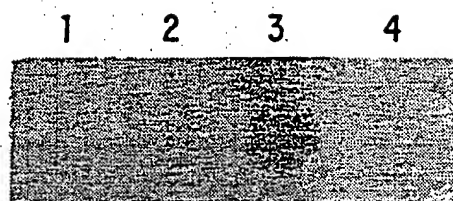


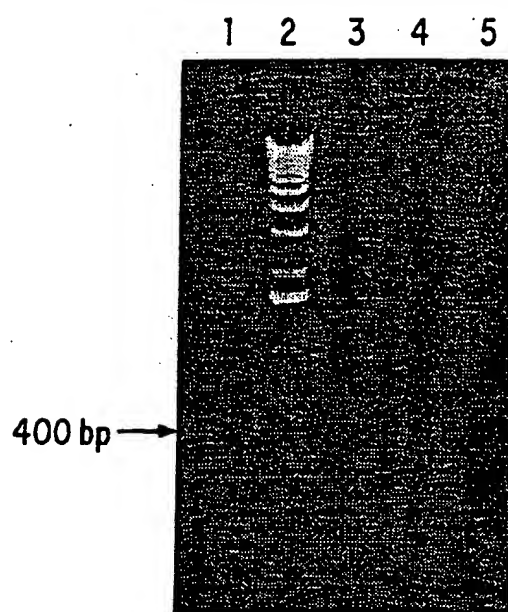
FIG. 3E

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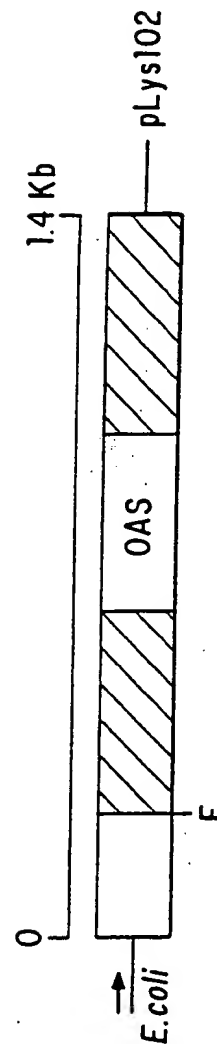
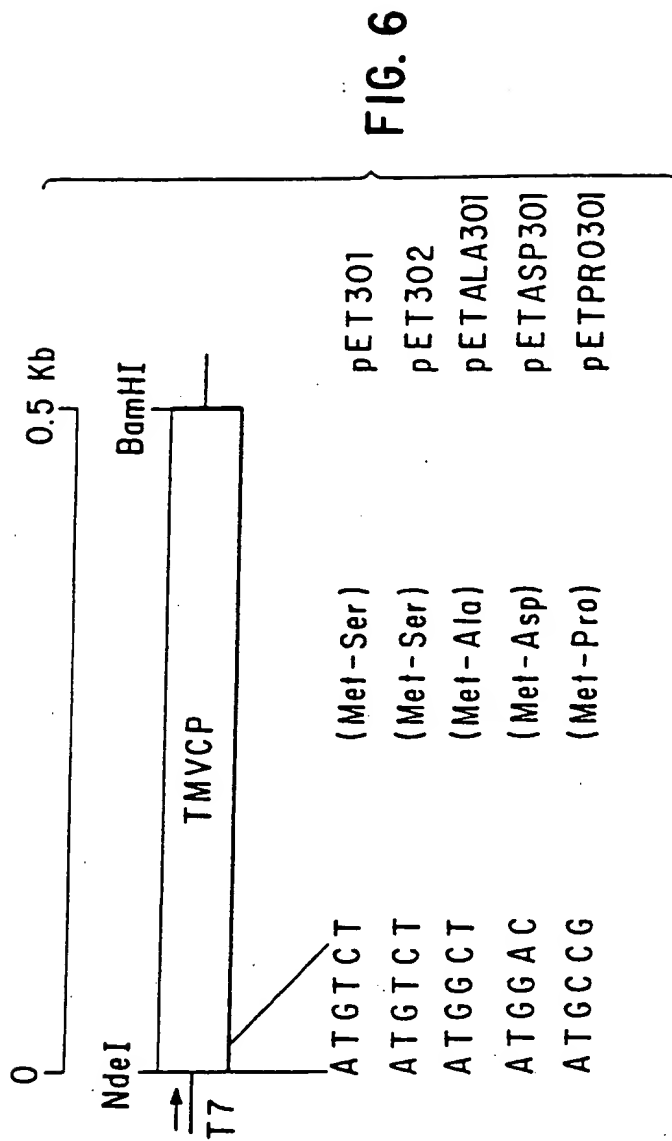


**FIG. 4**

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**FIG. 5**

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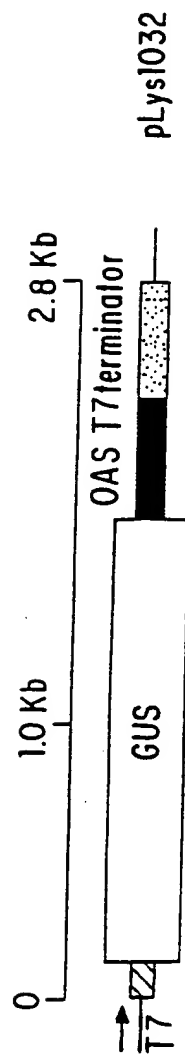


FIG. 8

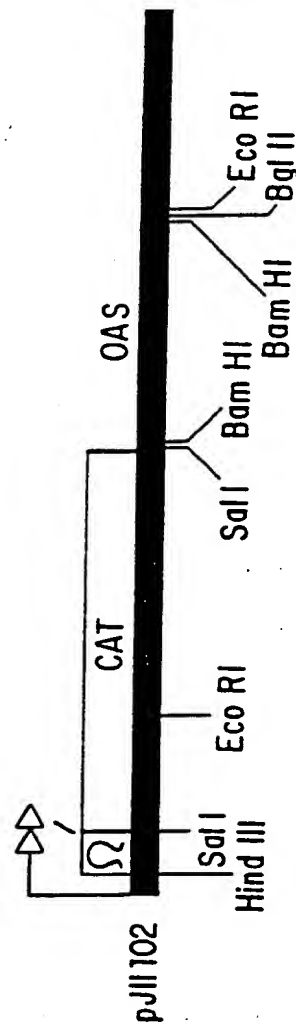


FIG. 9

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5118  
 |  
 GGUGGUCACG GGCAGUGGA ACUUGCCUGA CAAUUGCAGA GGAGGUGUGA GCGUGUGUCU 60  
 GGUGGACAAA AGGAUGGAAA GAGCCGACGA GCCCACUCUC GGAUCUUACU ACACAGCAGC 120  
 UGCAAAGAAA AGAUUUCAGU UCAAGGUCGU UCCCAAUUU GCUAUAACCA CCCAGGACGC 180  
 GAUGAAAAAC GUCUGGCAAG UUUUAGUUAA UAUUAGAAAU GUGAAGAUGU CAGCGGGUUU 240  
 CUGUCCGCUU UCUCUGGAGU UUGUGUCGGU GUGUAUUGUU UAUAGAAUA AUUAAAAUU 300  
 AGGUUUGAGA GAGAAGAUUA CAAACGUGAG AGACGGAGGG CCCAUGGAAC UUACAGAAGA 360  
 AGUCGUUGAU GAGUUCAUGG AAGAUGUCCC UAUGUCGAUC AGGCUUGCAA AGUUUCGAUC 420  
 UCGAACCGGA AAA 433  
 |  
 5550

FIG.10

UGAGAGACGG AGGGCCCAUG GAACUUACAG AAGAAGUCGU UGAUGAGUUC AUGGAAGAUG 60  
 UCCCUAUGUC GAUCA 75

FIG.11

SUBSTITUTE SHEET (RULE 26)



## INTERNATIONAL SEARCH REPORT

 International application No.  
 PCT/US93/10396

## A. CLASSIFICATION OF SUBJECT MATTER

IPC(S) : C12P 19/34; C12N 1/21, 7/01, 15/00; B65D 71/00

US CL : 435/91.32, 172.3, 235.1, 252.3, 252.33; 206/223

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 435/317.1, 810; 935/16

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

APS, Biosis, Derwent World Patents Index. Keywords: tobacco mosaic, TMV, coli, coat, assemb?, his, bacteri?, pseudovir?, recombinant.

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	Nucleic Acids Research, volume 16, number 8, issued 1988, Sleat et al, "Selective recovery of foreign gene transcripts as virus-like particles in TMV-infected transgenic tobaccos", pages 3127-3140, see entire document.	1-52
A	Bio/Technology, volume 4, issued July 1986, Haynes et al, "Development of a genetically-engineered candidate polio vaccine employing the self-assembling properties of the tobacco mosaic virus coat protein", pages 637-641, see entire document.	1-52

☒ Further documents are listed in the continuation of Box C.
 ☐ See patent family annex.

	Special categories of cited documents:	T	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
A	document defining the general state of the art which is not considered to be part of particular relevance		
E	earlier document published on or after the international filing date	X	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
L	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	Y	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
O	document referring to an oral disclosure, use, exhibition or other means		
P	document published prior to the international filing date but later than the priority date claimed	G	document member of the same patent family

Date of the actual completion of the international search

04 December 1993

Date of mailing of the international search report

JAN 31 1994

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## INTERNATIONAL SEARCH REPORT

International application No.  
PCT/US93/10396

## C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	Biochemistry, volume 29, issued 1990, Shire et al, "Preparation and properties of recombinant DNA derived tobacco mosaic virus coat protein", pages 5119-5126, see entire document.	1-52
A	Nucleic Acids Research, volume 17, number 2, issued 1989, Jupin et al, "Direct recovery of <i>in vitro</i> transcripts in a protected form suitable for prolonged storage and shipment at ambient temperatures", page 815, see entire document.	1-52
A	Journal of General Virology, volume 72, issued 1991, Jagadish et al, "Expression of potyvirus coat protein in <i>Escherichia coli</i> and yeast and its assembly into virus-like particles", pages 1543-1550, see entire document.	1-52